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## Ecological impacts of the austral-most population of *Crassostrea gigas* in South America: a matter of time?

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**Abstract** The Pacific oyster *Crassostrea gigas* is one of the most invasive species worldwide. This oyster has a preponderant ecological role in the invaded environments, for example structuring the benthic community through the provision of micro-habitats. Twenty-five years after its introduction in Argentina, the species is colonizing new areas along the coast, extending northwards and southwards its local distribution. In this study, we provide the first ecological characterization of the southern-most population of *C. gigas*; where the composition, density, richness and diversity of the macroinvertebrate assemblages associated with zones with oysters were compared with zones where it is absent at four different times of the year. Additionally, the main epibionts taxa settled on the oyster shells were studied. Our results showed differences in the assemblage composition between zones. However, these differences were not consistent throughout the year. Furthermore, density, richness and diversity were higher in the zones with

oysters only in one of the surveys and the parameters did not differ between zones in the remaining months. Moreover, the majority of oysters were used as settlement substrate by the sessile common species present in the area. Thus, our work provides new information about the ecology of *C. gigas* in recently invaded areas that enhance our understanding of the role that facilitation plays in physically stressful ecosystems and the importance that density and time since the invasion may have in the engineering effects of the species.

**Keywords** *Crassostrea gigas* · Invasive species · Facilitation · Epibiosis · Patagonia

### Introduction

Invasive species affect natural ecosystems in several ways. For example, changing ecosystem processes (Carlton 1999; Hoffmeister et al. 2005), generating a loss of biodiversity (Vitousek et al. 1997; Sala et al. 2000), introducing diseases and parasites (Hallegraeff 1998; Torchin et al. 2002), altering habitats (Schwindt et al. 2001; Crooks 2002) and modifying community functioning (Byers 1999; Grosholz et al. 2000). Specifically, facilitation by invasive species plays an important role in physically stressful ecosystems (Crooks 2002; Sousa et al. 2009). Many marine invasive plants (Bruno 2000; Brusati and Grosholz 2006; Neira et al. 2006), macroalgae (Strong et al. 2006; Byers et al. 2010; Irigoyen et al. 2011) and invertebrate species (Schwindt et al. 2001; Sousa et al. 2009; Sellheim et al. 2010) have profound architectural consequences on the ecosystem structure where they arrive. Examples from the literature suggest that introduced ecosystem engineers that increase habitat complexity or heterogeneity tend to change the abundances and/or species richness (Crooks 2002). Furthermore, several introduced species not only facilitate the survival of native species but also support the establishment and spread of other invaders (Simberloff and Von Holle 1999; Simberloff 2006). Thus,

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**Fig. 1** Map showing the different localities invaded by the Japanese oyster *Crassostrea gigas* on the coast of Argentina

while invasive species are considered among the top-five threats to native biodiversity (Vitousek et al. 1997; Sala et al. 2000), their arrival in the environments could lead to an increase in local richness.

Oyster introductions probably began as early as the seventeenth century, when the so-called Portuguese oyster (*Crassostrea angulata*) arrived in Europe from Asia (Carlton 1999). Overall, oysters have been introduced and established permanently in at least 30 countries outside their native ranges (Carlton 1999; Ruesink et al. 2005; Padilla 2010). The Pacific oyster *Crassostrea gigas*, a species native to Japan, is currently the most widely bivalve under aquaculture around the world and has been extensively introduced for cultivation (Orensanz et al. 2002; Ruesink et al. 2005; Padilla 2010). The Pacific oyster continues invading shores around the world, and is currently an invader in most of the temperate zones and some tropical areas worldwide (Ruesink et al. 2005; Padilla 2010). Introduced oysters have innumerable ecological impacts and do not only simply replace native oyster species (Ruesink et al. 2005). Oysters are considered autogenic engineer species because they change the physical structure, complexity and heterogeneity of the environment through their own structure (Jones et al. 1997; Ruesink et al. 2005; Sousa et al. 2009). The species is also an allogenic engineer since can decrease turbidity, increase light penetration in the water column and affect near bed flows and sedimentation processes (Ruesink et al. 2005). By suspension feeding, they influence energy flow and nutrient cycling at the scale of entire estuaries. Not only invasive oysters behave as engineers species; the eastern oyster *Crassostrea virginica* also exhibits the ability to form reefs and generate different effects on local communities in North America, where this species is native (Meyer and Townsend 2000; Shervette and Gelwick 2008). The transformations generated by oysters, both native and invasive, affect the associated species composition and their relative abundances since the interstices between oysters provide refuge for other species, and shells can

be used as a substratum for the settlement of algae and invertebrates (Meyer and Townsend 2000; Ruesink et al. 2005).

In South America, *Crassostrea gigas* was introduced to Chile and Perú in the Pacific Ocean, and to Brazil and Argentina in the Atlantic (Orensanz et al. 2002). In Argentina, its introduction dates back to 1982 when oysters from Coquimbo (Chile) were transplanted to Anegada Bay (~40°S, Fig. 1) with cultivation purposes (Orensanz et al. 2002). After a few months, the culture was abandoned and the oysters were left in the field (Orensanz et al. 2002). Fifteen years later, the settlement of oysters increased explosively and densities reached up to 120 recruits  $m^{-2}$  in the same area where they were introduced (Orensanz et al. 2002). After this main introduction, installations of other cultivation areas in the coast of Argentina were authorized between 1998 and 1999 (Orensanz et al. 2002). Currently, only a small population of *C. gigas* has been detected in the marshes of Samborombón Bay (~36°S, Fig. 1) possibly derived from one of these new cultivation areas (Giberto et al. 2012, but see Lomovasky et al. 2014). Since 2010, *C. gigas* has expanded its range to the north (~38°S, dos Santos and Fiori 2010, Fig. 1) and south (~41°S, Roche et al. 2010, Fig. 1) of the main introduction area. Thus, the invasion by this species could be reaching a new peak of expansion after a lag time in its invasion process (Crooks 2005). Although some ecological studies were performed (Escapa et al. 2004; Giberto et al. 2012; Croce and Parodi 2012), little is known about the effects of this species on native communities and, as often occurs, there is no information of how this species behaves in recently invaded areas. Indeed, this study is the first to investigate the potential impacts of *C. gigas* at the beginning of the establishment of a new population. The aim of the study is to provide the first ecological characterization of the southern-most population of *C. gigas* and to discuss possible facilitation processes of local fauna associated with its recent invasion. To do this, we compare the composition, density, richness and diversity

of the macroinvertebrate assemblages in zones with and without *C. gigas* and we repeat this in four times throughout one year. The main taxa of epibionts on *C. gigas* and their cover are also provided. Considering that density of a population is one of the key factors scaling ecosystem engineering impacts, we hypothesize that the effect generated by the presence of oysters in recent invaded environments will be less evident than that recorded in areas where the species has spent several decades established and populations reached higher densities.

## Materials and methods

### Study site

The study was performed in a fine sandstone rocky shore of the central Argentinean Patagonia, El Cóndor (41°03'S, 62°49'W, Fig. 1) where the tidal regime is semidiurnal with a mean amplitude of 4 m. The region is characteristic of the arid steppe, with predominant winds from the southwest, annual precipitation of 200 mm and mean annual temperature of 12 °C (Paruelo et al. 1998). In this rocky shore, *Crassostrea gigas* was recently found forming the austral-most population since its first introduction in the southwestern Atlantic coast with mean density of 0.11 ind m<sup>-2</sup> (Roche et al. 2010). The distribution of the oysters in the shore, mostly solitary individuals, is concentrated in the middle intertidal where the mytilid *Brachidontes rodriguezii* and the barnacles *Amphibalanus improvisus* and *Balanus glandula* are the dominant species.

### *Crassostrea gigas* as habitat-forming species

In order to compare the composition, density, richness and diversity of macroinvertebrates in areas with and without *C. gigas* (hereafter with oyster and without oyster zones) macrofaunal samples were collected. Fourteen samples per zone (10 cm × 10 cm) were obtained randomly at the middle intertidal of the rocky shore. Surveys were performed at four different times of the year: November 2009, April 2010, September 2010 and February 2011 to see if the effect of oysters was consistent throughout the year. In the field, the samples were collected separated by at least one meter, and interspersed in the middle intertidal and throughout the entire study sites. Each benthic sample was scraped off the surface, bagged, labelled and stored. In the laboratory, samples were sieved using a 0.5 mm mesh. The organisms retained on the mesh were fixed in 4 % formalin and preserved in 70 % ethanol. All organisms were identified to the lowest taxonomic level possible under a dissecting stereo microscope (80×), using updated taxonomic keys and the invertebrate reference collection from the CENPAT (<http://www.cenpat-conicet.gob.ar/>). A voucher of the taxa collected was deposited in the General Invertebrate Collection (CNP-INV). Moreover, we requested

the assistance of taxonomic specialists for some particular taxa. Afterwards, total density (individuals 100 cm<sup>-2</sup>), richness and Shannon diversity (Shannon and Weaver 1949) were calculated for each sample.

To determine if there were significant differences in invertebrate community composition between zones and months data were statistically analyzed using a permutational analysis of variance (PERMANOVA) with Primer 6 software (Primer-E, Plymouth; v.6.1.7) (Anderson et al. 2008). PERMANOVA compares the F statistics to a distribution generated by multiple random permutations of the analyzed data, thus liberating it from the formal assumptions of traditional ANOVA (Anderson 2001; Anderson et al. 2008). PERMANOVA model was performed considering zones (with and without) as a fixed factor and sampling months (4 levels) as a random factor (9999 permutations). Pairwise post hoc tests were performed for each month to identify where the differences occurred. The abundance of all invertebrate species was square-root transformed in order to down-weight the most abundant species. A similarity percentage analysis (SIMPER) was used to determine the taxa responsible for the differences between groups. To visualise multivariate patterns in benthic assemblages, non-metric multidimensional scaling (nMDS) was used as an ordination method. PERMANOVA was performed using a Bray–Curtis similarity matrix with a dummy variable. The PERMANOVA routine creates a non-parametric, permutational analogue of ANOVA when applied to univariate data (Anderson 2001; Anderson et al. 2008). Thereby, PERMANOVA models were also used to determine if there were significant differences in density, richness and diversity (9999 permutations). Pairwise post hoc tests were performed for each month to identify differences between zones for the three variables.

### Fouling on *Crassostrea gigas*

Epibionts on *C. gigas* were registered in each of the four sampling months (n = 14 oysters per month, except for April n = 10). The oyster size (maximum length × maximum width) was measured to the nearest mm with digital calliper (precision ± 0.01). The epibiotic organisms settled on the oyster shells were identified to the lowest taxonomic level possible under a dissecting stereo microscope (80×) and then the percentage cover of sessile organisms was estimated. Mobile organisms identified on the shells were also registered and counted. Differences in the percentage cover among oysters collected in different months were evaluated with a one way fixed ANOVA (Zar 1999). Normality and homogeneity of variance assumptions were evaluated with Kolmogorov–Smirnov and Levene tests, respectively. Parametric correlation analysis was performed to evaluate the relationship between the percentage cover and the size of the oyster (Zar 1999).

**Table 1** Mean density of invertebrate taxa (ind 100 cm<sup>-2</sup>) in zones with and without oysters

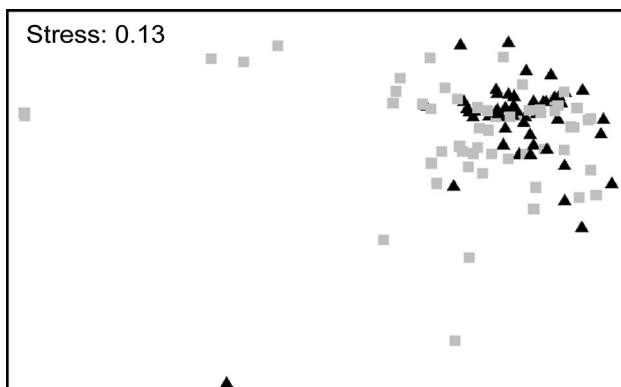
Phylum	Principal taxonomic group	Taxa	Zone	
			With oysters	Without oysters
Annelida	Polychaeta	Nereididae	0.20	0.05
		Syllidae	1.20	2.16
		Capitellidae	0.29	2.23
		Orbiniidae	0.13	0.23
		Lumbrineridae	0.07	0.04
Mollusca	Bivalvia	Spionidae		0.11
		Mytilidae	178.34	236.32
		Gasteropoda		
Arthropoda	Decapoda	<i>Costoanachis sertulariarum</i>	0.13	0.04
		<i>Parvanachis paessleri</i>		0.02
	Isopoda	<i>Cyrtograpsus altimanus</i>	0.91	0.20
		<i>Edotia</i> cf. <i>transversa</i>	0.02	0.05
	Amphipoda	<i>Amphithoe valida</i>	0.39	0.25
		<i>Monocorophium insidiosum</i>	0.48	0.18
	Tanaidacea	<i>Tanais dulongii</i>	0.02	
		Cirripedia		
	Diptera	<i>Balanus glandula</i>	8.14	9.59
		<i>Amphibalanus improvisus</i>	8.55	3.96
	Picnogonida	Chironomidae	0.50	0.86
		<i>Achelia assimilis</i>		0.02
Cephalorhyncha	Priapulida	sp. not identified	0.02	0.02
Cnidaria	Hydrozoa	<i>Amphisbetia operculata</i>	×	×
		<i>Eudendrium ramosum</i>	×	×
		<i>Plumularia setacea</i>	×	×
		<i>Ectopleura crocea</i>	×	×
		<i>Bicrisia edwardsiana</i>	×	×

The species *Aulacomya atra*, *Brachidontes rodriguezii* and *Mytilus* sp. were grouped together as Mytilidae since many individuals were juveniles in which the correct species identification was not possible. Presence was registered for Hydrozoa and Bryozoa

## Results

### *Crassostrea gigas* as habitat-forming species

A total of 27 macroinvertebrate taxa were found in the surveys (Table 1). The most abundant taxa for both zones were the mytilids (*Aulacomya atra*, *Brachidontes rodriguezii* and *Mytilus* sp.). Multivariate analysis showed that the invertebrate assemblage of the zones with oysters was different from zones without oysters (Fig. 2; Table 2). In MDS representation, samples from the same zone were



**Fig. 2** nMDS based on Bray–Curtis similarities of square-root transformed abundances associated to the zones with oysters (grey squares) and without oysters (black triangles)

more closely related to each other (Fig. 2). Besides, samples from the zones with oysters presented an average similarity of 74 % and those from the oyster zone 60 %. The average dissimilarity between the two zones was close to 40 % (Table 2). SIMPER analysis also showed that the taxa responsible for these differences changed throughout the year, being impossible to determine which of the taxa characterized each zone (Table 2). For example, the exotic barnacle *Balanus glandula* was more abundant in with oyster zone in November and September but the opposite result was found for April and February (Table 2). Furthermore, mytilids were the taxa that contributed mostly to the observed differences in assemblage structure between zones, regardless of the zone considered (SIMPER, top 40 % for all the surveys; Table 2); thus they were excluded in the density comparisons and their abundances analyzed separately. Community composition, density, richness and diversity differed significantly between zones; PERMANOVA detected a significant interaction between the factors Zones and Months showing that the differences between zones were not consistent over time and changed during the course of the study (Table 3). Pairwise comparisons showed that assemblage composition significantly differed between zones in the 4 months (Table 4). Density, richness, and diversity, instead, were higher in with oyster zone than in without oyster zone in November (Fig. 3; Table 4). Density of mytilids differed significantly between zones (PERMANOVA: pseudo- $f_{\text{zone}} = 2.42$ ,  $P > 0.05$ ; pseudo- $f_{\text{month}} = 5.28$ , pseudo- $f_{\text{zone} \times \text{month}} = 3.47$ ,  $P < 0.05$ ),

**Table 2** SIMPER routine results showing the taxa that mostly contributed to dissimilarity between zones

Month	Av. diss.	With av. abund.	Without av. abund.	Taxa	Contrib. (%)	Cum. (%)
November	39.08	11.51	12.70	Mytilidae	41.43	41.43
		2.36	0.10	<i>Balanus glandula</i>	17.59	59.03
		2.21	0.07	<i>Amphibalanus improvisus</i>	14.05	73.08
		0.56	0.81	Syllidae	6.51	79.59
April	42.87	9.71	5.73	Mytilidae	47.94	47.94
		1.01	2.26	<i>Balanus glandula</i>	14.83	62.77
		0.59	1.79	Capitellidae	10.04	72.81
		1.02	0.65	<i>Amphibalanus improvisus</i>	7.94	80.75
September	42.77	10.25	15.16	Mytilidae	42.16	42.16
		0.99	1.71	<i>A. improvisus</i>	13.48	55.65
		1.57	0.23	<i>Balanus glandula</i>	9.33	64.98
		0.00	1.33	Capitellidae	8.91	73.89
February	28.41	0.56	0.94	Chironomidae	7.25	81.14
		17.71	16.04	Mytilidae	42.73	42.73
		0.60	1.31	<i>Balanus glandula</i>	13.14	55.86
		1.00	1.89	Syllidae	11.51	67.37
		1.32	0.40	<i>Amphibalanus improvisus</i>	11.21	78.58

Lists were truncated when cumulative percentage reached 75 %. Average abundances were calculated with square-root transformed data. *Av. diss.* average dissimilarity, *Av. abund.* average abundance, *Contrib. %* contribution percentage, *Cum. %* cumulative percentage

**Table 3** PERMANOVAs results (df: zone = 1, month = 3, zone × month = 3)

Source	Composition			Density		
	Pseudo-f	<i>P</i>	Perms	Pseudo-f	<i>P</i>	Perms
Zone	1.79	0.2095	425	1.74	0.2647	425
Month	3.57	0.0001	9911	3.32	0.0042	9943
Zone × month	3.84	0.0001	9910	3.82	0.0011	9943

Source	Richness			Diversity		
	Pseudo-f	<i>P</i>	Perms	Pseudo-f	<i>P</i>	Perms
Zone	0.08	0.8567	425	0.41	0.5610	425
Month	1.85	0.1156	9948	1.02	0.3877	9955
Zone × month	3.07	0.0138	9946	4.37	0.0068	9960

Since mytilids account for about 90 % of the total abundance in both zones, they were excluded in the density comparisons

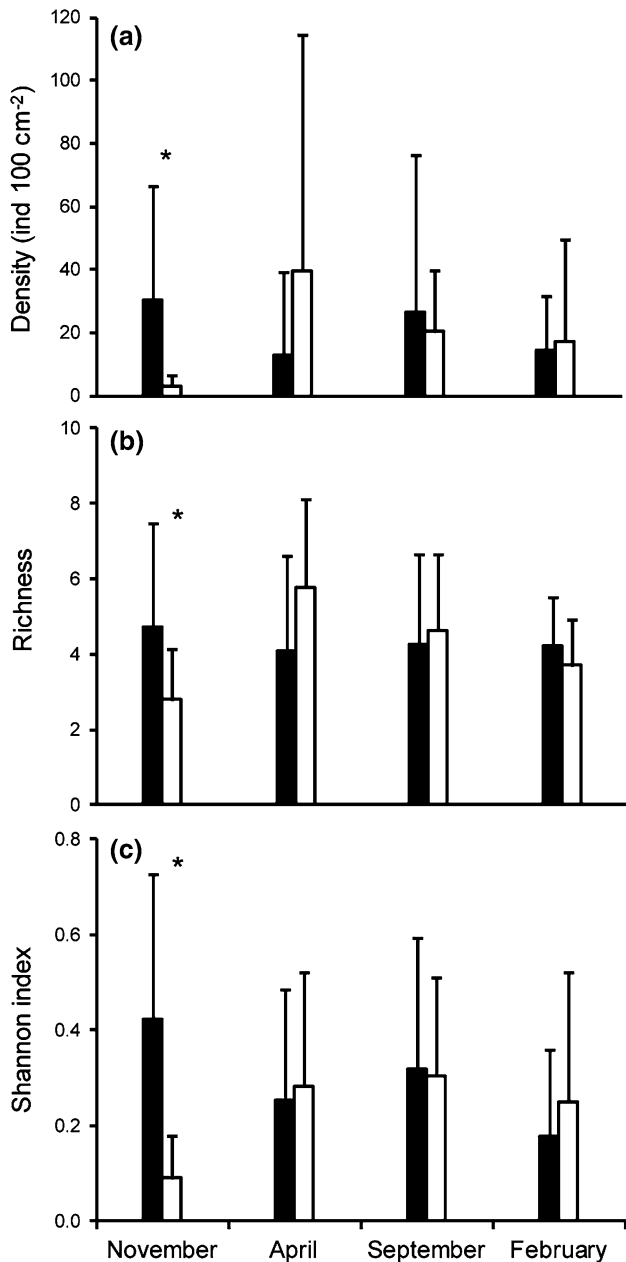
**Table 4** Results of the pairwise comparisons for the PERMANOVAs

	Composition			Density		
	<i>T</i>	<i>P</i>	Perms	<i>T</i>	<i>P</i>	Perms
November	1.96	0.004	9946	0.91	0.5504	9880
April	2.30	0.001	9936	2.71	0.0003	9925
September	2.39	0.007	9931	2.71	0.0030	9924
February	1.73	0.010	9941	1.07	0.2910	9934

	Richness			Diversity		
	<i>T</i>	<i>P</i>	Perms	<i>T</i>	<i>P</i>	Perms
November	1.94	0.0449	2905	4.06	0.0003	9904
April	1.76	0.0667	5522	0.37	0.7348	9917
September	0.56	0.6698	2718	0.15	0.9323	9909
February	0.91	0.3797	311	0.75	0.4568	9904





**Fig. 3** Mean ( $\pm$  SD) values of density (a), richness (b) and Shannon Diversity Index (c) of macroinvertebrates associated to the zones with oysters (black) and without oysters (white) for each month. Asterisks indicate significant differences between zones. Since mytilids account for about 90 % of the total abundance in both zones, they were excluded in the density comparisons

being higher in the without oyster zone in April and September (pairwise comparisons  $P < 0.05$ ).

#### Fouling on *Crassostrea gigas*

Over 90 % of the studied oysters showed epibiotic organisms. The size distribution of the oysters ranged from 10 to 60 cm<sup>2</sup>. Ten taxa of sessile organisms (cor-

responding to four species of bivalves, two barnacles, two hydrozoans and two bryozoans) and five of mobile ones (three polychaetes, one decapod and one insect larvae) were registered. The most abundant taxa were the mytilids and the barnacles *Balanus glandula* and *Amphibalanus improvisus*. The mean percentage cover was 34.8 (SD = 34.4). There were no significant differences in the percentage cover among months ( $F = 2.06$ ,  $df = 3$ ,  $P > 0.05$ ) and the size of the oysters was not significantly correlated with the epibionts percentage cover ( $P > 0.05$ ).

#### Discussion

After three decades since the first introduction of *Crassostrea gigas* in Argentina, the species expanded its range to two new locations, El C ndor (Roche et al. 2010, this study) the southern-most and Blanca Bay (dos Santos and Fiori 2010) the northern-most area. In our study, the invertebrate assemblage of the zones with oysters was different from zones without oysters. These differences in the community composition were not consistent throughout the year of study and the taxa that contributed to the identity of the zones also changed during the study. Density, richness, and diversity were higher in the oyster zone only in one of the surveys. In addition, oyster shells provide substratum for the settlement of several species, mostly mytilids and barnacles. Together, the results show that the presence of the invasive oyster *C. gigas* would not change markedly natural community patterns. However, the presence of oysters facilitated the establishment of some species probably by providing shelter and settlement sites. Since the invasion of oysters in the studied rocky shore is a recent phenomenon, the influence of the species in the environment could increase in the near future along with an increase in the population density. In this way, the potential effects of *C. gigas* could become more important if repeated recruitments occur and could resemble the effect of the species registered for other locations (Escapa et al. 2004; Padilla 2010).

Temporal dynamics in the invasion process, as well as the timing of many invasion-related events, are hard to predict (Crooks and Soul  1999; Simberloff 2003; Crooks 2005). For example, the time course of invasions often includes long initial periods of relative inactivity followed by drastic changes in invader dynamics (Crooks 2005). This phenomenon of lag times has become an increasingly recognized aspect of invasion biology (Crooks and Soul  1999; Simberloff 2003; Crooks 2005; Bortolus 2008). This kind of time lags can be found throughout the invasion process, including in the arrival, establishment, and in the impacts generated by invaders. For example, populations often grow exponentially in the early phases of invasion, and this gives rise to an inherent lag (Crooks 2005). Such is the case of *Crassostrea gigas* invasion in Argentina. After its

introduction in 1982, the species colonized the area at low densities and fifteen years later, settlement and densities increased explosively (Orensanz et al. 2002). Now, after 30 years the species has shown the capacity of expansion outside the original area of introduction, establishing populations to the north and the south (dos Santos and Fiori 2010 and Roche et al. 2010, respectively). Thereby, density of *C. gigas* in Argentina varies radically throughout its current introduced range (Fig. 1). In the first invaded areas (Anegada Bay) population reached average densities of 300 ind m<sup>-2</sup> (Borges 2006); however in new invaded areas (El C ndor) mean density is 0.11 ind m<sup>-2</sup> (Roche et al. 2010).

The density of a population is one of the key factors scaling ecosystem engineering impact; at low abundance, species are unlikely to cause community or ecosystem level change (Borthagaray and Carranza 2007; Padilla 2010; Harley and O'Riley 2011). In a detailed review of the engineering effect of *Crassostrea gigas*, Padilla (2010) highlighted the fact that the potential effects of oysters clearly depend on its population densities. For example, in rocky shores dominated by mussels (like our study site) oysters at low densities can produce an overgrowth of benthic species, generate local impacts on diversity and increase diversity of species sensitive to stress. However, at higher densities, oysters could also displace mussels, increase sedimentation, increase local diversity and alter species composition (Padilla 2010). Therefore, the major differences in density between early and newly invaded areas registered for Argentina could be explaining why the oysters in El C ndor did not modify sharply natural community patterns. In Anegada Bay, the first invaded intertidal in Argentina, the dense population of *C. gigas* has led to the creation of shallow intertidal reefs (Borges 2006). In this system, the abundance of several invertebrate species is highest within the reefs created by the oysters and, consequently, the density and rate of foraging of shorebirds is greatest in these areas, probably due to an increase in preys supply (Escapa et al. 2004). Also, Anegada Bay reefs provide microhabitats for macroalgal settlement (Croce and Parodi 2012). In Samboromb n Bay, oysters are grouped forming small reefs that, unlike Anegada Bay, are made up by small sizes individuals (shell height up to 37 mm, Giberto et al. 2012). Crustaceans, bivalves and other invertebrates are associated to these reefs (Giberto et al. 2012). Given the low density of oysters, reef habitats are still absent in our study site and could explain the differences in the effect generated by oysters found between Anegada Bay or Samboromb n Bay and El C ndor. Nevertheless, oysters at El C ndor offer a new settlement site for other organisms, such as mussels and barnacles, which in turn create micro-habitats for other benthic organisms (Prado and Castilla 2006; Sellheim et al. 2010; Sueiro et al. 2011). In this manner, differences in the time of establishment of the populations of *C. gigas* in Argentina have generated a density gradient between different invaded sites. This gradient, allow us to understand the complex relationship between the density of oysters and the effect exerted on communities, resulting

in greater impacts in places where oyster populations are denser and longer time since invasion.

It should be noted that some other environmental variation between Anegada Bay and El C ndor, such as the exposure of the coast or bottom type, could be determining *Crassostrea gigas* behavior and should be considered and analyzed in future research. In this sense, how oysters would affect the benthic community in the new invaded area of El C ndor is difficult to predict. In addition, the effect generated by oysters may vary depending on the presence of other engineering species (Padilla 2010). In general, the extremely high stress imposed by weather conditions in Patagonian rocky shores results in the dominance of habitat-forming species, mainly coralline algal turfs and mussel beds (Liuzzi and L pez Gappa 2008; Silliman et al. 2011). Mussels were dominant in our samples and we found differences in mussel densities between with oyster and without oyster zones in April and September. However, the increments in mussel densities were not consistent correlated with changes in the abundances of the other members of the invertebrate assemblage, and neither would influence the results found. Regardless of this, further investigation should focus on the potential interactive effects of oysters and mussels. Moreover, it is known that abiotic and biotic factors jointly contribute to the risk of proliferation of invasive species (Case 1990; Alpert et al. 2000; Ruesink 2007). In this respect, during the field surveys we observed that fine sediment load was considerably high (up to 500 g 100 cm<sup>-2</sup>). Since oysters are filter-feeders, the suspension of sediment could be affecting, indirectly and negatively, the feeding activity and therefore their survival. Alternatively, density of oyster's larvae can be reduced due to predation by filter-feeders (Pechenik et al. 2004; Porri et al. 2008; Troost 2010). In fact, oysters in particular, appear to consume large numbers of larvae, including those of their own species (Pechenik et al. 2004; Troost 2010). Predation by filter-feeders present in the area, as mytilids and the own *C. gigas*, could be an important source of variation in settlement success of oysters and possibly will affect population density. Additional studies are needed to evaluate how the different factors are conditioning the establishment and proliferation of the new population of *C. gigas*.

In densely populated marine environments like rocky shores, competition for space is critical (Connell 1961; Jackson 1977), thus the advantage of colonizing new surfaces is probably the major benefit of epibiosis (Wahl 1989). In El C ndor, more than 90 % of *Crassostrea gigas* individuals presented incrustated organisms mainly composed by sessile taxa. Among the most abundant organisms settled on oyster shells, we found the barnacles *Balanus glandula* and *Amphibalanus improvisus*. This association between the presence of exotic oysters and the barnacles deserves special attention since *B. glandula* is an invasive species and *A. improvisus* a cryptogenic one for Argentina (Orensanz et al. 2002). In the same way, it has been observed that the invasive filamentous

algae *Polysiphonia morrowii* (Raffo et al. 2014) recruits on *C. gigas* reefs in Anegada Bay (Croce and Parodi 2012). Both cases illustrate how invasive species are able to facilitate the colonization of other exotic species (Simberloff and Von Holle 1999; Simberloff 2006). As a result, ecological consequences of this kind of facilitation process could be severe (Simberloff and Von Holle 1999; Simberloff 2006; Sellheim et al. 2010) and should be studied in detail. Furthermore, several advantages and disadvantages may occur as a result of epibiosis, for both substrata organism and epibiont species (Wahl 1989, 2008). In the case of *C. gigas*, the settlement of large epibionts may increase the risk of dislodgement, reduce growth, increase the susceptibility to predation or affect feeding activities of oysters (Buschbaum and Reise 1999; Thielges 2005a; da Gama et al. 2008). On the other hand, water-retaining epibionts can slow down oyster desiccation or even camouflage them from predators (Thielges 2005b; Ramsby et al. 2012). Thereby, performing experimental studies of the potential effects of epibiosis could be useful to have a complete scenario of the impact of *C. gigas* on the communities.

To summarize, in the last years *Crassostrea gigas* started a new period of expansion in its invasion in Argentina. Our study suggests that the recent arrival of the invasive oyster in El Cóndor has not yet changed intensely the benthic community. Given the engineering activities commonly exhibited by the species, possible habitat alteration may occur in the short term if the density of the oyster increases. Taking into account that the population density in El Cóndor is still low, temporal and spatial monitoring becomes critical and allows this system to be an excellent opportunity to study the performance of an invader and its progressive effects. Such studies will help to focus concerns about how invaders face abiotic and/or biotic constraints. Environmental managers, decision makers and government authorities should exploit and benefit from the early detection of this invasion expansion as to prevent the spread of biological invasions and monitor their effects.

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